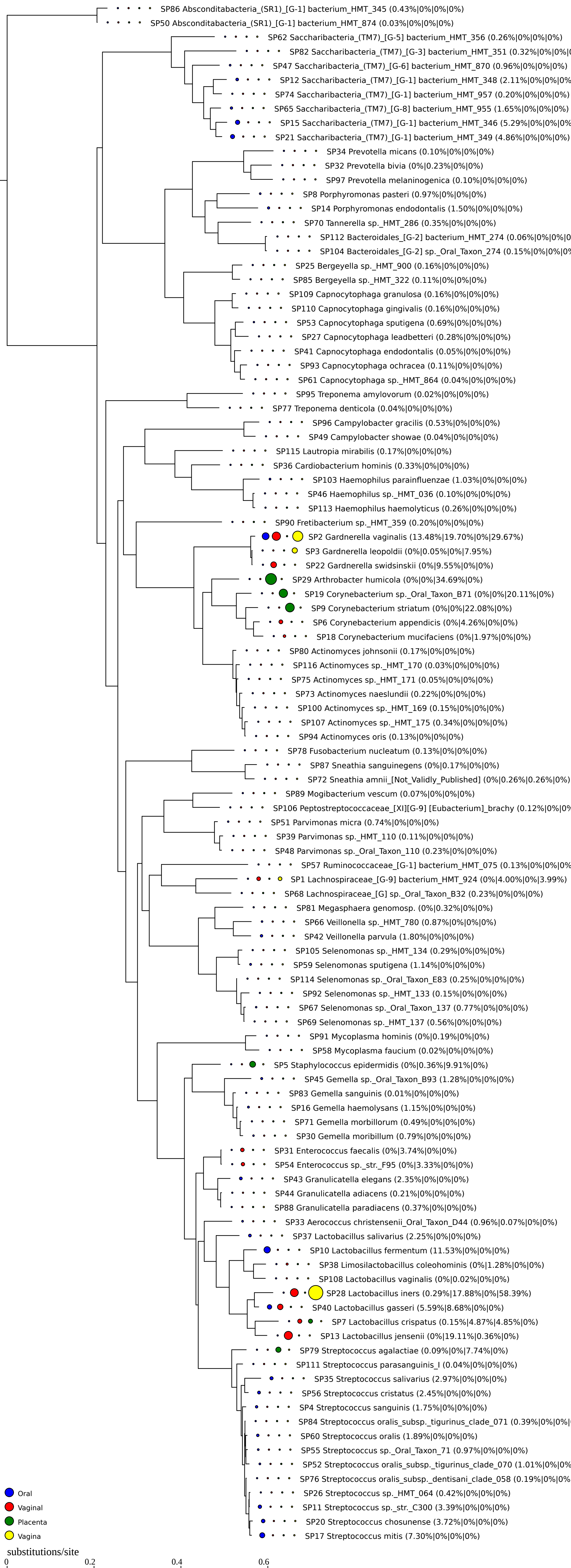
SP86 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0.43%|0%|0%|0%)
SP50 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874 (0.03%|0%|0%|0%)
SP62 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356 (0.26%|0%|0%|0%)
SP82 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 (0.32%|0%|0%|
SP47 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0.96%|0%|0%|0%)
SP12 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (2.11%|0%|0%|0%
SP74 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957 (0.20%|0%|0%|0%)
SP65 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955 (1.65%|0%|0%|0%)
SP15 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (5.29%|0%|0%|0%
SP21 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 (4.86%|0%|0%|0%)
SP34 Prevotella micans (0.10%|0%|0%|0%)
SP32 Prevotella bivia (0%|0.23%|0%|0%)
SP97 Prevotella melaninogenica (0.10%|0%|0%|0%)
SP8 Porphyromonas pasteri (0.97%|0%|0%|0%)
SP14 Porphyromonas endodontalis (1.50%|0%|0%|0%)
SP70 Tannerella sp._HMT_286 (0.35%|0%|0%|0%)
SP112 Bacteroidales_[G-2] bacterium_HMT_274 (0.06%|0%|0%|0
SP104 Bacteroidales_[G-2] sp._Oral_Taxon_274 (0.15%|0%|0%|0%
SP25 Bergeyella sp._HMT_900 (0.16%|0%|0%|0%)
SP85 Bergeyella sp._HMT_322 (0.11%|0%|0%|0%)
SP109 Capnocytophaga granulosa (0.16%|0%|0%|0%)
SP110 Capnocytophaga gingivalis (0.16%|0%|0%|0%)
SP53 Capnocytophaga sputigena (0.69%|0%|0%|0%)
SP27 Capnocytophaga leadbetteri (0.28%|0%|0%|0%)
SP41 Capnocytophaga endodontalis (0.05%|0%|0%|0%)
SP93 Capnocytophaga ochracea (0.11%|0%|0%|0%)
SP61 Capnocytophaga sp._HMT_864 (0.04%|0%|0%|0%)
SP95 Treponema amylovorum (0.02%|0%|0%|0%)
SP77 Treponema denticola (0.04%|0%|0%|0%)
SP96 Campylobacter gracilis (0.53%|0%|0%|0%)
SP49 Campylobacter showae (0.04%|0%|0%|0%)
SP115 Lautropia mirabilis (0.17%|0%|0%|0%)
SP36 Cardiobacterium hominis (0.33%|0%|0%|0%)
SP103 Haemophilus parainfluenzae (1.03%|0%|0%|0%)
SP46 Haemophilus sp._HMT_036 (0.10%|0%|0%|0%)
SP113 Haemophilus haemolyticus (0.26%|0%|0%|0%)
SP90 Fretibacterium sp._HMT_359 (0.20%|0%|0%|0%)
SP2 Gardnerella vaginalis (13.48%|19.70%|0%|29.67%)
SP3 Gardnerella leopoldii (0%|0.05%|0%|7.95%)
SP22 Gardnerella swidsinskii (0%|9.55%|0%|0%)
SP29 Arthrobacter humicola (0%|0%|34.69%|0%)
SP19 Corynebacterium sp._Oral_Taxon_B71 (0%|0%|20.11%|0%)
SP9 Corynebacterium striatum (0%|0%|22.08%|0%)
SP6 Corynebacterium appendicis (0%|4.26%|0%|0%)
SP18 Corynebacterium mucifaciens (0%|1.97%|0%|0%)
SP80 Actinomyces johnsonii (0.17%|0%|0%|0%)
SP116 Actinomyces sp._HMT_170 (0.03%|0%|0%|0%)
SP75 Actinomyces sp._HMT_171 (0.05%|0%|0%|0%)
SP73 Actinomyces naeslundii (0.22%|0%|0%|0%)
SP100 Actinomyces sp._HMT_169 (0.15%|0%|0%|0%)
SP107 Actinomyces sp._HMT_175 (0.34%|0%|0%|0%)
SP94 Actinomyces oris (0.13%|0%|0%|0%)
SP78 Fusobacterium nucleatum (0.13%|0%|0%|0%)
SP87 Sneathia sanguinegens (0%|0.17%|0%|0%)
SP72 Sneathia amnii_[Not_Validly_Published] (0%|0.26%|0.26%|0%)
SP89 Mogibacterium vescum (0.07%|0%|0%|0%)
SP106 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_brachy (0.12%|0%|0%
SP51 Parvimonas micra (0.74%|0%|0%|0%)
SP39 Parvimonas sp._HMT_110 (0.11%|0%|0%|0%)
SP48 Parvimonas sp._Oral_Taxon_110 (0.23%|0%|0%|0%)
SP57 Ruminococcaceae_[G-1] bacterium_HMT_075 (0.13%|0%|0%|0%
SP1 Lachnospiraceae_[G-9] bacterium_HMT_924 (0%|4.00%|0%|3.99%)
SP68 Lachnospiraceae_[G] sp._Oral_Taxon_B32 (0.23%|0%|0%|0%)
SP81 Megasphaera genomosp. (0%|0.32%|0%|0%)
SP66 Veillonella sp._HMT_780 (0.87%|0%|0%|0%)
SP42 Veillonella parvula (1.80%|0%|0%|0%)
SP105 Selenomonas sp._HMT_134 (0.29%|0%|0%|0%)
SP59 Selenomonas sputigena (1.14%|0%|0%|0%)
SP114 Selenomonas sp._Oral_Taxon_E83 (0.25%|0%|0%|0%)
SP92 Selenomonas sp._HMT_133 (0.15%|0%|0%|0%)
SP67 Selenomonas sp._Oral_Taxon_137 (0.77%|0%|0%|0%)
SP69 Selenomonas sp._HMT_137 (0.56%|0%|0%|0%)
SP91 Mycoplasma hominis (0%|0.19%|0%|0%)
SP58 Mycoplasma faucium (0.02%|0%|0%|0%)
SP5 Staphylococcus epidermidis (0%|0.36%|9.91%|0%)
SP45 Gemella sp._Oral_Taxon_B93 (1.28%|0%|0%|0%)
SP83 Gemella sanguinis (0.01%|0%|0%|0%)
SP16 Gemella haemolysans (1.15%|0%|0%|0%)
SP71 Gemella morbillorum (0.49%|0%|0%|0%)
SP30 Gemella moribillum (0.79%|0%|0%|0%)
SP31 Enterococcus faecalis (0%|3.74%|0%|0%)
SP54 Enterococcus sp._str._F95 (0%|3.33%|0%|0%)
SP43 Granulicatella elegans (2.35%|0%|0%|0%)
SP44 Granulicatella adiacens (0.21%|0%|0%|0%)
SP88 Granulicatella paradiacens (0.37%|0%|0%|0%)
SP33 Aerococcus christensenii_Oral_Taxon_D44 (0.96%|0.07%|0%|0%)
SP37 Lactobacillus salivarius (2.25%|0%|0%|0%)
SP10 Lactobacillus fermentum (11.53%|0%|0%|0%)
SP38 Limosilactobacillus coleohominis (0%|1.28%|0%|0%)
SP108 Lactobacillus vaginalis (0%|0.02%|0%|0%)
SP28 Lactobacillus iners (0.29%|17.88%|0%|58.39%)
SP40 Lactobacillus gasseri (5.59%|8.68%|0%|0%)
SP7 Lactobacillus crispatus (0.15%|4.87%|4.85%|0%)
SP13 Lactobacillus jensenii (0%|19.11%|0.36%|0%)
SP79 Streptococcus agalactiae (0.09%|0%|7.74%|0%)
SP111 Streptococcus parasanguinis_I (0.04%|0%|0%|0%)
SP35 Streptococcus salivarius (2.97%|0%|0%|0%)
SP56 Streptococcus cristatus (2.45%|0%|0%|0%)
SP4 Streptococcus sanguinis (1.75%|0%|0%|0%)
SP84 Streptococcus oralis_subsp._tigurinus_clade_071 (0.39%|0%|0%|
SP60 Streptococcus oralis (1.89%|0%|0%|0%)
SP55 Streptococcus sp._Oral_Taxon_71 (0.97%|0%|0%|0%)
SP52 Streptococcus oralis_subsp._tigurinus_clade_070 (1.01%|0%|0%
SP76 Streptococcus oralis_subsp._dentisani_clade_058 (0.19%|0%|0%
SP26 Streptococcus sp._HMT_064 (0.42%|0%|0%|0%)
SP11 Streptococcus sp._str._C300 (3.39%|0%|0%|0%)
SP20 Streptococcus chosunense (3.72%|0%|0%|0%)
SP17 Streptococcus mitis (7.30%|0%|0%|0%)
Oral
Vaginal
Placenta
Vagina
substitutions/site
0
0.2
0.4
0.6